40. The process of claim 31 wherein the DNA probe corresponds to the nucleotide sequence coding for glycoproteins gp41 or gp120 of the HIV-3 retrovirus or the complement thereof.--

REMARKS

The active claims in this case are claims 31 and 37-40.

The specification has been amended to recite the relationship with the parent cases, and to incorporate portions of the preliminary amendment filed August 23, 1999 in parent case 09/379,270. A marked up version of the amendments to the Specification is attached hereto. The original application was filed without an abstract of the disclosure. This Preliminary Amendment is being filed to provide such an abstract. The title has been amended to reflect the subject of the present claims.

The specification at page 3 has been amended to introduce the current classification, HIV-1 subtype O virus, for what had been termed "HIV-3" in the original application. After the inventors first reported on their discovery of HIV-3, specifically variant ANT₇₀, the medical and scientific community recognized that HIV-3 should more appropriately be classified as a subtype of HIV-1. This subtype was designated "O" where O stands for "outliers". Several journal articles have been provided in the preceding case Serial No. 08/486,836 to substantiate the scientific recognition that HIV-3, e.g., ANT₇₀, is now classified as HIV-1 subtype O.

The new claims find support at page 30, line 25 through page 32, line16; pages 37-40; pages 50-53; and original claims 25-31. A marked up version of the claim amendments is attached.

It is believed that no fee is due; however, should any fees under 37 C.F.R. §§ 1.16 to 1.21 be required for any reason, the Commissioner is authorized to deduct said fees from Deposit Account No. 01-2508/11362.0025.DVUS03.

Respectfully submitted,

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Date:

May 9, 2001

MARKED-UP VERSION OF CLAIMS

31. (Amended) A process for the detection of HIV-3 retrovirus or of its RNA in a biological liquid or tissue, characterized by contacting nucleic acids contained in said biological liquid or tissue with a <u>DNA</u> probe containing [a nucleic acid according to any of claims 25 to 30] at least 360 contiguous sequences corresponding to the genomic RNA of HIV-3 retrovirus under stringent hybridization conditions, washing the hybrid formed with a solution preserving said stringent conditions, and detecting the hybrid formed.

The process of claim 31 wherein the DNA probe is: --37.

37. The process of claim 31 wherein the DIVA process							
_	10	20 TGAAGATACA	30 CATAAAGAAA	40 TACTGATGTG	GAAGTTTGAT	AGATCTCTAG	
	70	80 TGTTGCTATG	90 ATAACTCACC	100 CAGAGCTCTT	110 CCAGAAGGAC	120 TAAAAACTGC	
	130	140 ATTGCTGACA	150 CTGTGGAACT	160 TTCCAGCAAA	170 GACTGCTGAC	180 ACTGCGGGGA	
	190	200 GGAGGGACAG	210 GGGGCGGTTC	220 GGGGAGTGGC	230 TAACCCTCAG	240 AAGCTGCATA	
	250	260 CTTTCTGCTT	270 GTACCGGGTC	280 TCGGTTAGAG	290 GACCAGGTCT	300 GAGCCCGGGA	
TAAGCAGC	310	320 CTCTAGCTGA	330 ACCCGCTCGT	340 TAACGCTCAA	350 TAAAGCTTGC	360 CTTGAGTGAG	
A							

or the complement thereof.

The process of claim 31 wherein the DNA probe is: 38.

38. The proc	60				
10	20	30 AAAAGGTAAA	40 TTTGAGGGAT	50 GGGCAGCAGT	AAGAGAAAGA
AACATGGGAA	80	90	100	110	120
	CTAGAACTTT	CCCTGAGTCT	GAACCATGCG	CACCTGGAGT	AGGACAGATC
ATGAGAAGAA	140	150	160	170	180
	TAGCAGCTAG	AGGAGGGATA	CCAAGTTCCC	ATACTCCTCA	AAACAATGCA
TCCAGGGAAT 190 GCCCTTGCAT	200	210	220	230	240
	TCCTAGAAAG	TCACCAAGAG	GAAGAAGTAG	GTTTTCCAGT	AGCACCTCAA
250	260	270	280	290	300
GTGCCTCTAA	GGCCAATGAC	CTATAAAGGA	GCATTTGACC	TCAGCTTCTT	TTTAAAAGAA

310 320 330 340 350 360 AAGGGAGGAC TGGAAGGGTT AATTTACTCC CATAAAAGAG CAGAAATCCT GGATCTTTGG GTGTATAA or the complement thereof.

- 39. The process of claim 31 wherein the DNA probe corresponds to the nucleotide sequence coding for proteins p12, p16 or p25 of the HIV-3 retrovirus or the complement thereof.
- 40. The process of claim 31 wherein the DNA probe corresponds to the nucleotide sequence coding for glycoproteins gp41 or gp120 of the HIV-3 retrovirus or the complement thereof.--